Abstract

A method for the isolation of CDRs in a de-
fined framework that is stable and soluble in reducing
5   environment is described as well as thus obtainable scFv.
Starting from such scFv with defined framework a scFv li-
brary can be generated wherein the framework is conserved
while at least one complementary determining region (CDR)
is randomized. Such library, e.g. in yeast cells, is
10   suitable for screening for antibody/CDR-interactions or
for screening for antibodies.